## RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 09/6/3./70Source: 160/6Date Processed by STIC: 5/9/05

## ENTERED



IFW16

RAW SEQUENCE LISTING DATE: 05/09/2005
PATENT APPLICATION: US/09/613,170 TIME: 16:20:18

Input Set : N:\Crf3\RULE60\09613170.raw
Output Set: N:\CRF4\05092005\I613170.raw

```
SEQUENCE LISTING
        (1) GENERAL INFORMATION:
             (i) APPLICANT: Ueda, Tetsufumi
                             Ozkan, Eric D.
      6
            (ii) TITLE OF INVENTION: Compositions And Methods For The
      8
                                      Inhibition Of Neurotransmitter Uptake Of Synaptic
      9
Vesicles
           (iii) NUMBER OF SEQUENCES: 6
     11
            (iv) CORRESPONDENCE ADDRESS:
     13
                   (A) ADDRESSEE: Medlen & Carroll, LLP
     14
                   (B) STREET: 220 Montgomery Street, Suite 2200
     15
     16
                   (C) CITY: San Francisco
                   (D) STATE: California
     17
                   (E) COUNTRY: United States of America
     18
                   (F) ZIP: 94104
     19
     21
             (v) COMPUTER READABLE FORM:
                   (A) MEDIUM TYPE: Floppy disk
     22
                   (B) COMPUTER: IBM PC compatible
     23
                   (C) OPERATING SYSTEM: PC-DOS/MS-DOS
     24
                   (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
     25
            (vi) CURRENT APPLICATION DATA:
     27
C--> 28
                   (A) APPLICATION NUMBER: US/09/613,170
C-->29
                   (B) FILING DATE: 10-Jul-2000
                   (C) CLASSIFICATION: 514
     30
           (vii) PRIOR APPLICATION DATA:
     32
                   (A) APPLICATION NUMBER: US/08/840,006
     33
                   (B) FILING DATE: 15-APR-1997
     34
          (viii) ATTORNEY/AGENT INFORMATION:
     36
     37
                   (A) NAME: Carroll, Peter G.
                   (B) REGISTRATION NUMBER: 32,837
     38
                   (C) REFERENCE/DOCKET NUMBER: UM-02786
     39
            (ix) TELECOMMUNICATION INFORMATION:
     41
                   (A) TELEPHONE: (415) 705-8410
     42
                   (B) TELEFAX: (415) 397-8338
     43
        (2) INFORMATION FOR SEQ ID NO: 1:
     46
             (i) SEQUENCE CHARACTERISTICS:
     48
                   (A) LENGTH: 10 amino acids
     49
                   (B) TYPE: amino acid
     50
                   (C) STRANDEDNESS: Not Relevant
     51
                   (D) TOPOLOGY: linear
     52
            (ii) MOLECULE TYPE: peptide
     54
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
     59
             Glu Ala Ala Leu Thr Ser Glu Glu Val Gly
     61
     62
                                                   10
             1
```

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PATENT APPLICATION: US/09/613,170 TIME: 16:20:18

Input Set: N:\Crf3\RULE60\09613170.raw
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```
64 (2) INFORMATION FOR SEQ ID NO: 2:
        (i) SEQUENCE CHARACTERISTICS:
66
             (A) LENGTH: 5 amino acids
67
             (B) TYPE: amino acid
68
             (C) STRANDEDNESS: Not Relevant
69
             (D) TOPOLOGY: linear
70
       (ii) MOLECULE TYPE: peptide
72
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
77
        Tyr His Arg Phe Lys
79
                         5
80
        1
82
       INFORMATION FOR SEQ ID NO: 3:
        (i) SEQUENCE CHARACTERISTICS:
84
             (A) LENGTH: 10 amino acids
85
             (B) TYPE: amino acid
86
             (C) STRANDEDNESS: Not Relevant
87
             (D) TOPOLOGY: linear
88
       (ii) MOLECULE TYPE: peptide
90
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
95
        Tyr His Arg Phe Lys Glu Leu Ser Thr Leu
97
98
                                              10
        INFORMATION FOR SEQ ID NO: 4:
100
         (i) SEQUENCE CHARACTERISTICS:
102
               (A) LENGTH: 20 amino acids
103
               (B) TYPE: amino acid
104
               (C) STRANDEDNESS: Not Relevant
105
               (D) TOPOLOGY: linear
106
108
        (ii) MOLECULE TYPE: peptide
113
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
         Tyr His Arg Phe Lys Glu Leu Ser Thr Leu Arg Arg Gln Lys Leu Glu
115
116
                                               10
                                                                    15
118
         Asp Ser Tyr Arg
119
                      20
        INFORMATION FOR SEQ ID NO: 5:
121 (2)
123
         (i) SEQUENCE CHARACTERISTICS:
124
               (A) LENGTH: 1200 amino acids
125
              (B) TYPE: amino acid
126
               (C) STRANDEDNESS: Not Relevant
               (D) TOPOLOGY: linear
127
129
        (ii) MOLECULE TYPE: peptide
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
134
         Met Asp Pro Ser Gly Val Lys Val Leu Glu Thr Ala Glu Asp Ile Gln
136
137
         Glu Arg Arg Gln Gln Val Leu Asp Arg Tyr His Arg Phe Lys Glu Leu
139
140
         Ser Thr Leu Arg Arg Gln Lys Leu Glu Asp Ser Tyr Arg Phe Gln Phe
142
143
                  35
         Phe Gln Arg Asp Ala Glu Glu Leu Glu Lys Trp Ile Gln Glu Lys Leu
145
             50
                                                       60
146
                                  55
         Gln Ile Ala Ser Asp Glu Asn Tyr Lys Asp Pro Thr Asn Leu Gln Gly
148
```

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/613,170

DATE: 05/09/2005

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Input Set : N:\Crf3\RULE60\09613170.raw
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149	65					70					75					80
151	Lys	Leu	Gln	Lys	His	Gln	Ala	Phe	Glu	Ala	Glu	Val	Gln	Ala	Asn	Ser
152					85					90					95	
154	Gly	Ala	Ile	Val	Lys	Leu	Asp	Glu	Thr	Gly	Asn	Leu	Met	Ile	Ser	Glu
155				100					105					110		
157	Gly	His	Phe	Ala	Ser	Glu	Thr	Ile	Arg	Thr	Arg	Leu	Met	Glu	Leu	His
158	_		115					120					125			
160	Arg	Gln	Trp	Glu	Leu	Leu	Leu	Glu	Lys	Met	Arg	Glu	Lys	Gly	Ile	Lys
161		130	_				135					140				
163	Leu	Leu	Gln	Ala	Gln	Asn	Leu	Val	Gln	Tyr	Leu	Arg	Glu	Cys	Glu	Asp
164	145					150					155					160
166	Val	Met	Asp	Trp	Ile	Asn	Asp	Lys	Glu	Ala	Ile	Val	Thr	Ser	Glu	Glu
167			_	-	165					170					175	
169	Leu	Gly	Gln	Asp	Leu	Glu	His	Val	Glu	Val	Leu	Gln	Lys	Lys	Phe	Glu
170				180					185					190		
172	Glu	Phe	Gln	Thr	Asp	Met	Ala	Ala	His	Glu	Glu	Arg	Val	Asn	Glu	Val
173			195					200					205			
175	Asn	Gln	Phe	Ala	Ala	Lys	Leu	Ile	Gln	Glu	Gln	His	Pro	Glu	Glu	Glu
176		210					215					220				
178	Leu	Ile	Lys	Thr	Lys	Gln	Asp	Glu	Val	Asn	Ala	Ala	Trp	Gln	Arg	Leu
179	225					230					235					240
181	Lys	Gly	Leu	Ala	Leu	Gln	Arg	Gln	Gly	Lys	Leu	Phe	Gly	Ala	Ala	Glu
182					245					250					255	
184	Val	Gln	Arg	Phe	Asn	Arg	Asp	Val	Asp	Glu	Thr	Ile	Ser	Trp	Ile	Lys
185				260					265					270		
187	Glu	Lys	Glu	Gln	Leu	Met	Ala	Ser	Asp	Asp	Phe	Gly	Arg	Asp	Leu	Ala
188			275					280					285			
190	Ser	Val	Gln	Ala	Leu	Leu	Arg	Lys	His	Glu	Gly	Leu	Glu	Arg	Asp	Leu
191		290					295					300				
193	Ala	Ala	Leu	Glu	Asp	Lys	Val	Lys	Ala	Leu	Cys	Ala	Glu	Ala	Asp	Arg
194	305					310					315					320
196	Leu	Gln	Gln	Ser	His	Pro	Leu	Ser	Ala	Thr	Gln	Ile	Gln	Val		Arg
197					325			_		330		_		_	335	
199	Glu	Glu	Leu	Ile	Thr	Asn	Trp	Glu	Gln	Ile	Arg	Thr	Leu		Ala	Glu
200				340					345			_		350		_
202	Arg	His	Ala	Arg	Leu	Asn	Asp		Tyr	Arg	Leu	Gln	Arg	Phe	Leu	Ala
203			355			_		360	_	_	_		365	_	_	
205	Asp	Phe	Arg	Asp	Leu	Thr	Ser	Trp	Val	Thr	Glu		Lys	Ala	Leu	Ile
206		370				_	375			_		380	_	_		_
208		Ala	Asp	Glu	Leu	Ala	Ser	Asp	Val	Ala	Gly	Ala	Glu	Ala	Leu	
209	385			_	_	390		_		_	395	_		<b>-</b>		400
211	Asp	Arg	His	Gln		His	Lys	Gly	Glu		Asp	Ala	His	Glu	_	Ser
212	•	_	_		405		_		<b>-</b>	410		_			415	•
214	Phe	Lys	Ser		Asp	Glu	Ser	Gly		Ala	Leu	Leu	Ala		Gly	His
215	_	_ <b>-</b>	_	420		<del>-</del>	_	<b>-</b>	425	_		i	_	430	~- <b>-</b>	
217	Tyr	Ala		Asp	Glu	Val	Arg		Lys	Leu	Thr	Val	Leu	Ser	Glu	GIu
218	_		435	_	_		_	440		_	_	_	445	~-3	<b>~</b>	~- 7
220	Arg		Ala	Leu	Leu	Glu		Trp	Glu	Leu	Arg		Gln	GIn	Tyr	Glu
221		450					455					460				

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223		Cys	Met	Asp	Leu		Leu	Phe	Tyr	Arg	_	Thr	Glu	Gln	Val	_
224	465	(T)	16-4	<b>0</b>	<b>T</b>	470	<b>~1</b>	7 J -	Dh a	T	475	7	<b>a</b> 1	7 ~~	T 0	480
226	ASI	Trp	мес	ser	-	GIII	Glu	Ala	Pne		ьeu	ASII	GIU	Asp		Gry
227	7	Dh.	T	7	485	77_ T	<u>ما</u>	77.	T	490	Ť	T	11: ~	<b>~1</b>	495	Dho
229	Asp	Pne	Leu	_	Ser	vai	Glu	Ala		Leu	ьys	гуѕ	HIS		Asp	Pne
230	<b>~1</b>	T	C 0 22	500	C 0	77.	<b>~1</b> =	<b>~1</b>	505	T	T]_	mh	ח ד ת	510	7 ~~	~1
232	GIU	ьуѕ		ьеи	ser	Ala	Gln		GIU	ьуѕ	me	TIII		ьeu	Asp	GIU
233	Dho	ת דת	515	Tara	Ton	т1.	C15	520	λαν	Uic	Φτ.τ~	הוג	525	C111	7 cm	Wal
235	Pne	530	THE	тур	ьeu	116	Gln	ASII	ASII	urs	ıyı		Mec	GIU	Asp	vai
236 238	ת 1 ת		7 ~~	7~~	7 cm	ת ד ת	535 Leu	Lou	Cor	7 ~~	7 ~~	540	בות	Leu	Uic	Glu
239	545	1111	Arg	Arg	Asp	550	пеп	ьец	261	Arg	555	Maii	AIa	ъсц	птэ	560
241		λla	Mot	λνα	7 ra		Ala	Cln	Leu	בות		Car	Dhe	Wic	T.e.u	
241	Arg	MIG	MEC	Arg	565	AIG	MIG	GIII	пеп	570	Asp	SEL	FIIC	1115	575	GIII
244	Gln	Dhe	Dhe	Ara		Sar	Asp	Glu	T.211		Sar	ሞተካ	V = 1	Δen		Lve
245	GIII	Liic	FIIC	580	Asp	261	wsh	Giu	585	цуэ	Ser	пр	Val	590	GIU	цуз
247	Met	Lve	Thr		Thr	Δsn	Glu	Δla		Lvc	Δsn	Pro	Ser		T.eu	Gln
248	MCC	цу	595	mu	1111	пор	OIU	600	- 7 -	цуб	nop	110	605	11011	Leu	0111
250	Glv	Lvs		Gln	Lvs	His	Gln		Phe	Glu	Δla	Glu		Ser	Δla	Asn
251	Q-1	610	• • • • • • • • • • • • • • • • • • • •	0111	Lys	1110	615	1114	1	OI u	1114	620	200	JC1		
253	Gln		Ara	Tle	Asp	Ala	Leu	Glu	īvs	Ala	Glv		Lvs	Leu	Ile	Asp
254	625	001	5		TIOP	630	200	024	<b>-</b> 10		635	<b>U</b>	_1.			640
256		Asn	His	Tvr	Ala		Asp	Glu	Val	Ala		Ara	Met	Asn	Glu	
257				- 1 -	645	-7-	F		. •	650					655	. •
259	Ile	Ser	Leu	Trp		Lvs	Leu	Leu	Glu		Thr	Glu	Leu	Lvs		Ile
260				660	1	-7-			665					670	1	
262	Lys	Leu	Arq	Glu	Ala	Asn	Gln	Gln	Gln	Gln	Phe	Asn	Arq	Asn	Val	Glu
263	•		675					680					685			
265	Asp	Ile	Glu	Leu	Trp	Leu	Tyr	Glu	Val	Glu	Gly	His	Leu	Ala	Ser	Asp
266	•	690			-		695				-	700				_
268	Asp	Tyr	Gly	Lys	Asp	Leu	Thr	Asn	Val	Gln	Asn	Leu	Gln	Lys	Lys	His
269 <sup>°</sup>	705	_	-	-	_	710					715					720
271	Ala	Leu	Leu	Glu	Ala	Asp	Val	Ala	Ala	His	Gln	Asp	Arg	Ile	Asp	Gly
272					725	_				730					735	
274	Val	Thr	Ile	Gln	Ala	Arg	Gln	Phe	Gln	Asp	Ala	Gly	His	Phe	Asp	Ala
275				740					745					750		
277	Glu	Asn	Ile	Lys	Lys	Lys	Gln	Glu	Ala	Leu	Val	Ala	Arg	Tyr	Glu	Ala
278			755					760					765			
280	Leu	Lys	Glu	Pro	Met	Val	Ala	Arg	Lys	Gln	Lys	Leu	Ala	Asp	Ser	Leu
281		770	•				775					780				
283	Arg	Leu	Gln	Gln	Leu	Phe	Arg	Asp	Val	Glu	Asp	Glu	Glu	Thr	Trp	Ile
284	785					790					795					800
286	Arg	Glu	Lys	Glu	Pro	Ile	Ala	Ala	Ser	Thr	Asn	Arg	Gly	Lys	Asp	Leu
287					805					810					815	
289	Ile	Gly	Val	Gln	Asn	Leu	Leu	Lys	Lys	His	Gln	Ala	Leu	Gln	Ala	Glu
290				820					825					830		
292	Ile	Ala	Gly	His	Glu	Pro	Arg	Ile	Lys	Ala	Val	Thr	Gln	Lys	Gly	Asn
293			835					840					845			
295	Ala	Met	Val	Glu	Glu	Gly	His	Phe	Ala	Ala	Glu	Asp	Val	Lys	Ala	Lys

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Input Set : N:\Crf3\RULE60\09613170.raw
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296			850					855					860				
298		Len		Glu	T.en	Δsn	Gln		Trn	Glu	Δla	T.em		Ala	Lvs	Ala	Ser
299		865	1110	Olu	Lcu	11011	870	Lys	P	<b>01</b> u	1114	875			270		880
301			Ara	Arg	Gln	Asn		G111	Asn	Ser	Len		Ala	Gln	Gln	Tvr	
302		0111	m 9	my	OIII	885	Dea	Olu	nop	JCI	890	<b>0111</b>	*****	0411	01	895	1110
304		Δla	Acn	Ala	Δen		בומ	Glu	Sar	Trn		Δτα	Glu	Twe	Glu		Tle
305		ATO	Aph	AIQ	900	Giu	AIG	Giu	561	905	MEC	rra	Gru	цуs	910	110	110
303		Wa 3	<u>را ، ، ، </u>	Ser		λαn	ጥኒታዮ	Cly	Tarc		Glu	λcn	Car	λla		λla	T.e.u
		vai	Gry	915	1111	Ash	ıyı	Gry	920	ASP	GIU	тэр	Der	925	GIU	AIG	<u>neu</u>
308		T 011	T		uic	C1.,	ת דת	T ON		Sor	7 cm	T 011	Sor		ጥኒታን	Clv	Ser
310		ьеu	-	Lys	птэ	GIU	Ala	935	Mec	ser	Asp	Leu	940	MIA	тут	Gry	261
311		Com	930	Cln	- נע	T 011	7 ~~		C1 n	ת הות	C1 5	Cor		ሽድα	Cln	Cln	v-1
313			TTE	Gln	Ald	ьеи	_	GIU	GIII	ATA	GIII		Cys	Arg	GIII	GIII	960
314		945	Dree	mb	7 ~~	7 ~~	950	mb se	<b>~1</b>	T	<i>C</i> 1,,	955	17.7	T 011	ת ב ת	T 011	
316		Ala	PIO	Thr	Asp	_	GIU	THE	GIY	ьуѕ		ьeu	val	ьeu	Ala		ıyı
317		3	m	<b>~1</b>	<b>~1</b>	965	0	D	7	<b>~</b> 1	970	Mla sa	Mob	T	T	975	7 an
319		Asp	Tyr	Gln		гаг	ser	Pro	Arg		vai	THE	мес	гуѕ		GLY	Asp
320		<b>-</b> 1 -	<b>T</b>	ml	980	<b>T</b>	<b>3</b>	0	ml	985	T	7	Ш	<b></b>	990	17.m. 7	<b>a</b> 1
322		тте	Leu	Thr	Leu	ьeu	Asn	ser			ьуѕ	Asp	Trp			vai	GIU
323		** 1	<b>.</b>	995	<b>3</b>	<b>~1</b> .	<b>01</b>	<b>5</b> 1	1000		<b>3</b> 1.	<b>71</b> -	<b></b>	100!	_	<b>T</b>	T
325		val		Asp	Arg	GIN	GIY			Pro	Ala	Ala	_		ьys	ьуѕ	Leu
326		_	1010		<b>~</b> 1	<b>a</b>			5	<b>~1</b>	<b>3</b>	<b>7</b>	1020		~1	<b>71.</b>	G1
328		_		Ala	Gin	Ser			Arg	GIU	Asn			GIU	GIU	GIN	
329		102!			_	_	1030		~3	<b>+3</b>		1035		m\	•	<b>T</b> ].	1040
331		Ser	He	Ala	Leu	_		GIU	GIn	TTE	_		GIn	Thr	Arg		
332		_	~3		<b>~</b> 3	104	_	_	-	•	1050		<b>~1</b>	77 - J	<b>0</b> 1	1055	
334		Lys	GLu	Ala	_		Val	Ser	Leu	_		гуѕ	GIn	vaı			Leu
335		_		_	106	-	~ 7	_	~3	1069	_	_	-	<b>~</b> 3	107		<b>a</b> 1
337		Tyr	His	Ser		Leu	GIu	Leu	_		Lys	Arg	ьуѕ			ьeu	GIU
338		_	_	107!			1			0	_	~1	<b>.</b>	108		<b>T</b>	<b>01</b>
340		Lys		Cys	Lys	Lys	Phe			Pne	Arg	GIU			GIU	ьeu	Gin
341			109		_	~3	_	109			_	1	110		~1	**. 7	<b>~</b> 3
343			-	Ile	Asn	GIU	_		Ala	Ala	Leu			GIU	GIU	vaı	_
344		110		•	<b>~</b> 1	<b>~1</b>	1110		**7	<b>.</b>	<b>01</b>	111!		Dl	7	7	1120
346		Ala	Asp	Leu	GIU			GIU	vaı	Leu		_	гуѕ	Pne	Asp	_	
347		~3	_	_	•	112		•	<b>~1</b>	0	1130		<b>T</b>	3	<b>7</b> 1.	113!	
349		GIn	Lys	Asp		-	АТа	Asn	GIU		_	Leu	гàг	Asp			гÀг
350		<b>-</b>		<b>9</b>	114	-		_	~-3	114!	_			~7	115		<b>~</b> 3
352		Val	Ala	Glu	_	Leu	Glu	Ser			Leu	Met	Ala			Val	GIn
353				115!			<b>-</b>	·	116				_	116			1
355		Ala		Gln	GIn	GIn	Glu		-	GLy	Met	Met			Asp	GIu	Thr
356			117		_	_		117	_		_		118				•
358		_		Lys	Thr	Ala			Trp	Lys	Ser			Leu	Met	Val	
359	_	118					119					119!	5				1200
362	(2)	INFO															
364		(i)		UENC						_							
365			(A)	) LEI	NGTH	: 11	51 aı	mino	acio	ds							
366																	
367			(C)		RAND	EDNE	SS: 1	Not 1	Rele	vant							

VERIFICATION SUMMARY

 $\phi^{\alpha} = (g^{\alpha})^{-1} = (g^{\alpha})^{-1}$ 

DATE: 05/09/2005

PATENT APPLICATION: US/09/613,170

TIME: 16:20:19

Input Set : N:\Crf3\RULE60\09613170.raw
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L:28 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]

L:29 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]